



PC10970.ST25.txt
SEQUENCE LISTING

<110> Pfizer Inc.

Harland, Lee

<120> Novel Polypeptide

<130> PC10970AGLK

<150> GB0101739.1

<151> 2001-01-23

<150> US 60/267,341

<151> 2001-02-08

<160> 3

<170> PatentIn version 3.1

<210> 1

<211> 918

<212> DNA

<213> Homo sapiens

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ctcctggtga aaatgaacac ccggtcagt accaccatgg cggtcattaa cttggtggtg 180
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ctcacgttcc tattctatgt ggtgatcctg gtcaccagat acctcatctt cttcaagtgc 360
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aaggttgcac ttataacga aatcttcttg agtgtaacag caattagctg ctatgatttg 840
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<213> Homo sapiens

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Thr Pro His Leu Ile Ser Leu Tyr Phe Ile Val Leu Ile Gly Gly Leu
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Val Gly Val Ile Ser Ile Leu Phe Leu Leu Val Lys Met Asn Thr Arg
35           40           45

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Ser Val Thr Thr Met Ala Val Ile Asn Leu Val Val Val His Ser Val
50           55           60

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Phe Leu Leu Thr Val Pro Phe Arg Leu Thr Tyr Leu Ile Lys Lys Thr
65           70           75           80

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Trp Met Phe Gly Leu Pro Phe Cys Lys Phe Val Ser Ala Met Leu His
85           90           95

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Ile His Met Tyr Leu Thr Phe Leu Phe Tyr Val Val Ile Leu Val Thr
100          105          110

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Arg Tyr Leu Ile Phe Phe Lys Cys Lys Asp Lys Val Glu Phe Tyr Arg
115          120          125

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Lys Leu His Ala Val Ala Ala Ser Ala Gly Met Trp Thr Leu Val Ile
130 135 140

Val Ile Val Val Pro Leu Val Val Ser Arg Tyr Gly Ile His Glu Glu
145 150 155 160

Tyr Asn Glu Glu His Cys Phe Lys Phe His Lys Glu Leu Ala Tyr Thr
165 170 175

Tyr Val Lys Ile Ile Asn Tyr Met Ile Val Ile Phe Val Ile Ala Val
180 185 190

Ala Val Ile Leu Leu Val Phe Gln Val Phe Ile Ile Met Leu Met Val
195 200 205

Gln Lys Leu Arg His Ser Leu Leu Ser His Gln Glu Phe Trp Ala Gln
210 215 220

Leu Lys Asn Leu Phe Phe Ile Gly Val Ile Leu Val Cys Phe Leu Pro
225 230 235 240

Tyr Gln Phe Phe Arg Ile Tyr Tyr Leu Asn Val Val Thr His Ser Asn
245 250 255

Ala Cys Asn Ser Lys Val Ala Phe Tyr Asn Glu Ile Phe Leu Ser Val
260 265 270

Thr Ala Ile Ser Cys Tyr Asp Leu Leu Leu Phe Val Phe Gly Gly Ser
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Arg
305

<210> 3

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<212> PRT

<213> Homo sapiens

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 Ile Ser Val Val Gly Phe Phe Gly Asn Gly Phe Val Leu Tyr Val Leu
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 Ile Lys Thr Tyr His Lys Lys Ser Ala Phe Gln Val Tyr Met Ile Asn
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 65 70 75 80
 Val Tyr Tyr Val His Lys Gly Ile Trp Leu Phe Gly Asp Phe Leu Cys
 85 90 95
 Arg Leu Ser Thr Tyr Ala Leu Tyr Val Asn Leu Tyr Cys Ser Ile Phe
 100 105 110
 Phe Met Thr Ala Met Ser Phe Phe Arg Cys Ile Ala Ile Val Phe Pro
 115 120 125
 Val Gln Asn Ile Asn Leu Val Thr Gln Lys Lys Ala Arg Phe Val Cys
 130 135 140
 Val Gly Ile Trp Ile Phe Val Ile Leu Thr Ser Ser Pro Phe Leu Met
 145 150 155 160
 Ala Lys Pro Gln Lys Asp Glu Lys Asn Asn Thr Lys Cys Phe Glu Pro
 165 170 175
 Pro Gln Asp Asn Gln Thr Lys Asn His Val Leu Val Leu His Tyr Val
 180 185 190
 Ser Leu Phe Val Gly Phe Ile Ile Pro Phe Val Ile Ile Ile Val Cys
 195 200 205
 Tyr Thr Met Ile Ile Leu Thr Leu Leu Lys Lys Ser Met Lys Lys Asn
 210 215 220
 Leu Ser Ser His Lys Lys Ala Ile Gly Met Ile Met Val Val Thr Ala
 225 230 235 240
 Ala Phe Leu Val Ser Phe Met Pro Tyr His Ile Gln Arg Thr Ile His
 245 250 255
 Leu His Phe Leu His Asn Glu Thr Lys Pro Cys Asp Ser Val Leu Arg
 260 265 270

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Met Gln Lys Ser Val Val Ile Thr Leu Ser Leu Ala Ala Ser Asn Cys
 275 280 285

Cys Phe Asp Pro Leu Leu Tyr Phe Phe Ser Gly Gly Asn Phe Arg Lys
 290 295 300

Arg Leu Ser Thr Phe Arg Lys His Ser Leu Ser Ser Val Thr Tyr Val
 305 310 315 320

Pro Arg Lys Lys Ala Ser Leu Pro Glu Lys Gly Glu Glu Ile Cys Lys
 325 330 335

val